

<https://helda.helsinki.fi>

---

## Low-Frequency and Rare-Coding Variation Contributes to Multiple Sclerosis Risk

International Multiple Sclerosis Genetics Consortium

2020

---

International Multiple Sclerosis Genetics Consortium , Saarela , J & Theunissen , C E 2020 ,  
' Low-Frequency and Rare-Coding Variation Contributes to Multiple Sclerosis Risk ' , Cell ,  
vol. 180 , no. 2 , 403 . <https://doi.org/10.1016/j.cell.2020.01.002>

---

<http://hdl.handle.net/10138/325814>

<https://doi.org/10.1016/j.cell.2020.01.002>

---

cc\_by

publishedVersion

---

*Downloaded from Helda, University of Helsinki institutional repository.*

*This is an electronic reprint of the original article.*

*This reprint may differ from the original in pagination and typographic detail.*

*Please cite the original version.*

# Low-Frequency and Rare-Coding Variation Contributes to Multiple Sclerosis Risk

International Multiple Sclerosis Genetics Consortium\*

\*Correspondence: [chris.cotsapas@yale.edu](mailto:chris.cotsapas@yale.edu) (Chris Cotsapas)

<https://doi.org/10.1016/j.cell.2020.01.002>

(Cell 175, 1679–1687.e1–e7; November 29, 2018)

It has come to our attention that in preparing the final version of this article, the authors inadvertently misspelled the last name of author Charlotte E. Teunissen as “Charlotte E. Theunissen.” This error has been corrected in the article online.

In the Editorial Note (Cell 178, 262, June 27, 2019), the editors refer to the original version of the published manuscript. That version contained a misspelled name, and as that has now been corrected, we are updating the Editorial Note as well.

